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Sequence Listing was accepted.

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Reviewer: Durreshwar Anjum

Timestamp: [year=2011; month=12; day=20; hr=11; min=36; sec=40; ms=752;
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Application No: 10584424 Version No: 2.0

Input Set:**Output Set:**

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Finished: 2011-12-12 13:25:02.321
Elapsed: 0 hr(s) 0 min(s) 2 sec(s) 882 ms
Total Warnings: 68
Total Errors: 0
No. of SeqIDs Defined: 80
Actual SeqID Count: 80

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Input Set:

Output Set:

Started: 2011-12-12 13:24:59.439
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Total Warnings: 68
Total Errors: 0
No. of SeqIDs Defined: 80
Actual SeqID Count: 80

Error code

Error Description

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Shuttleworth, Gillian
Finnis, Christopher John Arthur

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<151> 2003-12-23

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<170> PatentIn version 3.5

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<213> Saccharomyces cerevisiae

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Tyr Lys Pro Leu Phe Thr Glu Leu Ala Lys Lys Asn Arg Gly Leu Met
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Asn Phe Val Ser Ile Asp Ala Arg Lys Phe Gly Arg His Ala Gly Asn
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340 345 350

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Glu Ile Phe Glu Asn Gln Asp Ser Ser Val Phe Gln Leu Val Gly Lys
370 375 380

Asn His Asp Glu Ile Val Asn Asp Pro Lys Lys Asp Val Leu Val Leu
385 390 395 400

Tyr Tyr Ala Pro Trp Cys Gly His Cys Lys Arg Leu Ala Pro Thr Tyr
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Gln Glu Leu Ala Asp Thr Tyr Ala Asn Ala Thr Ser Asp Val Leu Ile
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Ala Lys Leu Asp His Thr Glu Asn Asp Val Arg Gly Val Val Ile Glu
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Gly Tyr Pro Thr Ile Val Leu Tyr Pro Gly Gly Lys Lys Ser Glu Ser
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Val Val Tyr Gln Gly Ser Arg Ser Leu Asp Ser Leu Phe Asp Phe Ile
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Lys Glu Asn Gly His Phe Asp Val Asp Gly Lys Ala Leu Tyr Glu Glu
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Ile Gly Asp Ala Ala Lys Asn Gln Ala Ala Met Asn Pro Ser Asn Thr
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Val Phe Asp Ala Lys Arg Leu Ile Gly Arg Asn Phe Asn Asp Pro Glu
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Glu Thr Ala Gly Gly Val Met Thr Lys Leu Ile Pro Arg Asn Ser Thr
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Ile Ser Thr Lys Lys Phe Glu Ile Phe Ser Thr Tyr Ala Asp Asn Gln
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Pro Gly Val Leu Ile Gln Val Phe Glu Gly Glu Arg Ala Lys Thr Lys
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Asp Asn Asn Leu Leu Gly Lys Phe Glu Leu Ser Gly Ile Pro Pro Ala
450 455 460

Pro Arg Gly Val Pro Gln Ile Glu Val Thr Phe Asp Val Asp Ser Asn
465 470 475 480

Gly Ile Leu Asn Val Ser Ala Val Glu Lys Gly Thr Gly Lys Ser Asn
485 490 495

Lys Ile Thr Ile Thr Asn Asp Lys Gly Arg Leu Ser Lys Glu Asp Ile
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Glu Lys Met Val Ala Glu Ala Glu Lys Phe Lys Glu Glu Asp Glu Lys
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Glu Ser Gln Arg Ile Ala Ser Lys Asn Gln Leu Glu Ser Ile Ala Tyr
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Glu Leu Gln Asp Ile Ala Asn Pro Ile Met Ser Lys Leu Tyr Gln Ala
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